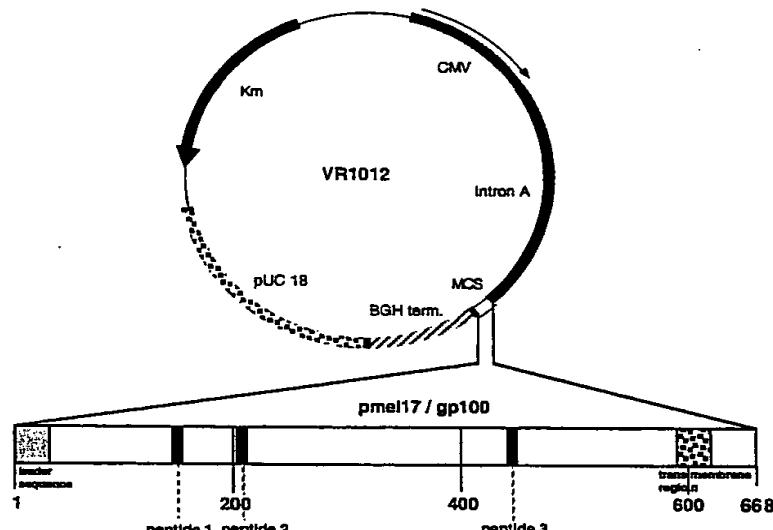


A**B**

		Peptide 1	Peptide 2	Peptide 3
human	1	MOLIVKRCOLLEAVICALLAVGATKVERNODWLGVSRQLETKAWNROLYPEWTEACRILDC		
mouse	1	MVCVORRSELFLYVVLISALLAVGALECSRQNQDWLGVPRQLTKTWNROLYPEWTEVGCSNC		
human	61	WRGGQVSLKVENDGPTLTGANASFIALNFPGSQKVLPDGCVIWWNNNTIINGSQVWGGQP		
mouse	61	WRGGQVSLRVENDGPTLVMGANASFIALHFPGSQKVLPDGCVIWWANNTIINGSQVWGGQP		
human	121	VYPOEADDACIFPDGGPCPSGSWSCKRSFVYVVKTWGQYHVQLGGPVSCLSTETCHAMLG		
mouse	121	VYPOEADDACIFPDGGPCPSGPKPEKRSFVYVVKTWGQYHVQLGGPVSRSSTATEHAKLG		
human	181	THTMEVTVYHRRGSRSYVPLAHSASSFTITDQVFPSVSVSQLALDCCNKHFLRNQPLTF		
mouse	181	THTMEVTVYHRRGSSYVPLAHSASSFTITDQVFPSVSVSQLALDCCNKHFLRNQPLTF		
human	241	AIQLHDPSGYLAEADLSYTWDFGDSSTGLISRAAVVTHTYLEPGPVTAQVVLQAAILPLAS		
mouse	241	AIQLHDPSGYLAEADLSYTWDFGDTIGTLISRAAVVTHTYLESGSVTAQVVLQAAILPLAS		
human	301	CGSSPVPGTTDGFPTAAEAPMNTAGCATTTEVVGTPGQPTAEPSCGTSVQVPTTEVIS		
mouse	301	CGSSPVPGTTDGFPTAAEAPCTSROGTTKVVGTPGQPTOPSGTTVQMPPTTEVTA		
human	361	TAFVQMATAESTGMPTEKVPSEVMCTTIAEMSITAEATGTEAEVSTIVVLSGTTAAQVMM		
mouse	361	TTSEOMIT-----SAVEDITLAEVSTIETGTTPTRP-----SGTTVAQMT		
human	421	TEWWETTARELPIPEPEGPDASLIVSTSITGSIICPLIDCTTIRLVKROVPLDCVLYRY		
mouse	403	TE-----GPDASLIPTOSSITGSIICPLIDDPDTIILVLRQVPLDCVLYRY		
human	481	GFSVLDIVQGIESAELQAVPSEGDAFELTVSCQGGLPKEACMEISSPGCQPPAORL		
mouse	448	GFSVLDIVQGIESAELQAVPSEGDAFELTVSCQGGLPKEACMDISSPGCQPPAORL		
human	541	CQPVLPSPFACQQLVLHQILKGGSPTYCLNVSLADANSLAVVSTOLIMPVPGILLTGQEAQI		
mouse	508	CQSVLPSPFACQQLVLHQILKGGSPTYCLNVSLADANSLAVVSTQLAVP-----GQDGGL		
human	601	GCVPLIVGILLVIMAVVIALSLIY----RRRLMKDFSYVQDPESSSEHWLRPRTIECSPI		
mouse	561	GCVPLIVGILLVIVAVVIALSLIYGIDLRSRAOFPKCENVAATAAPASCLRA-R---G-I		
human	657	GENSPLLSGQQV		
mouse	615	GENSPLLSGQQV		

Fig. 1

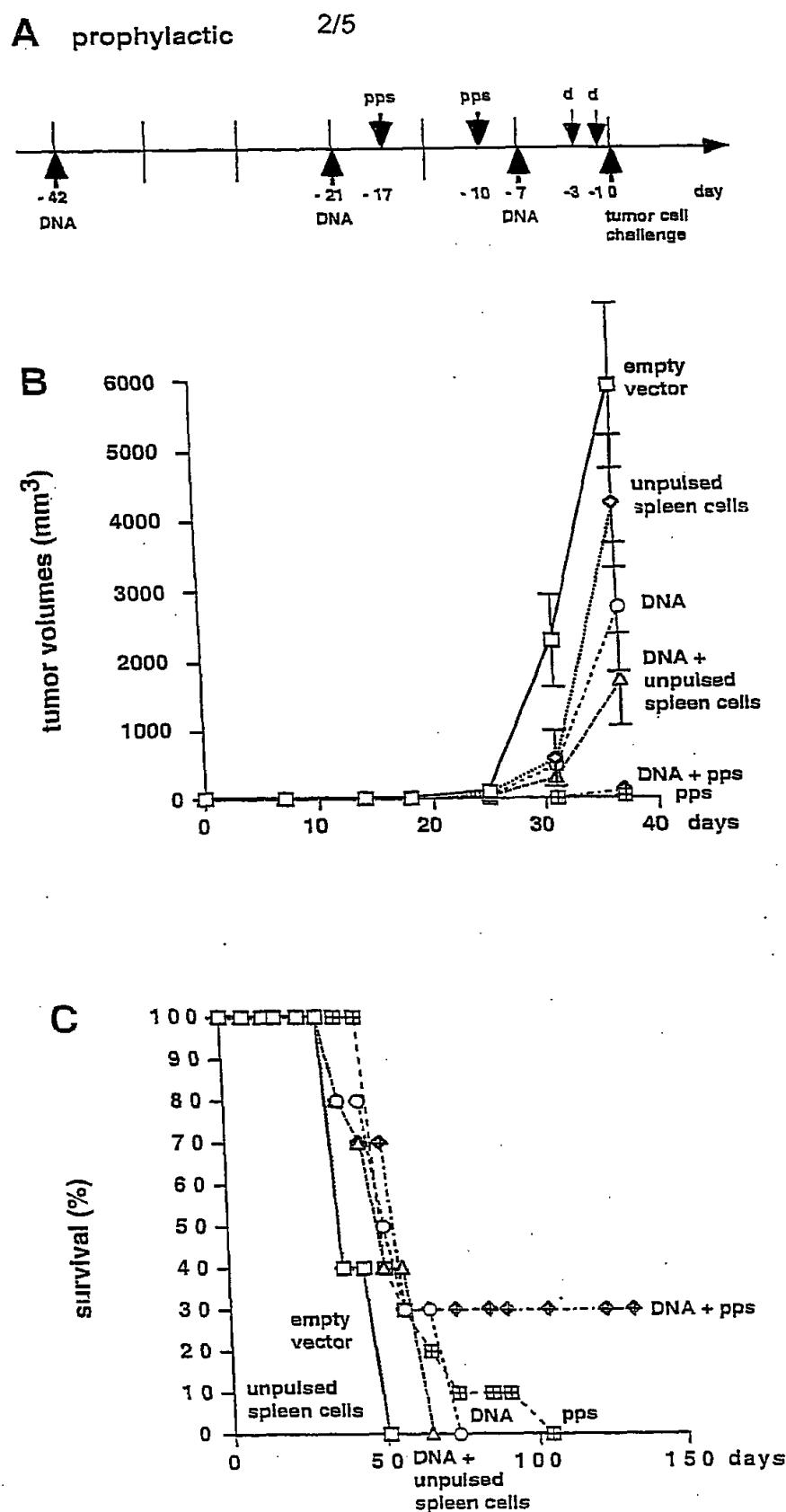


Fig. 2

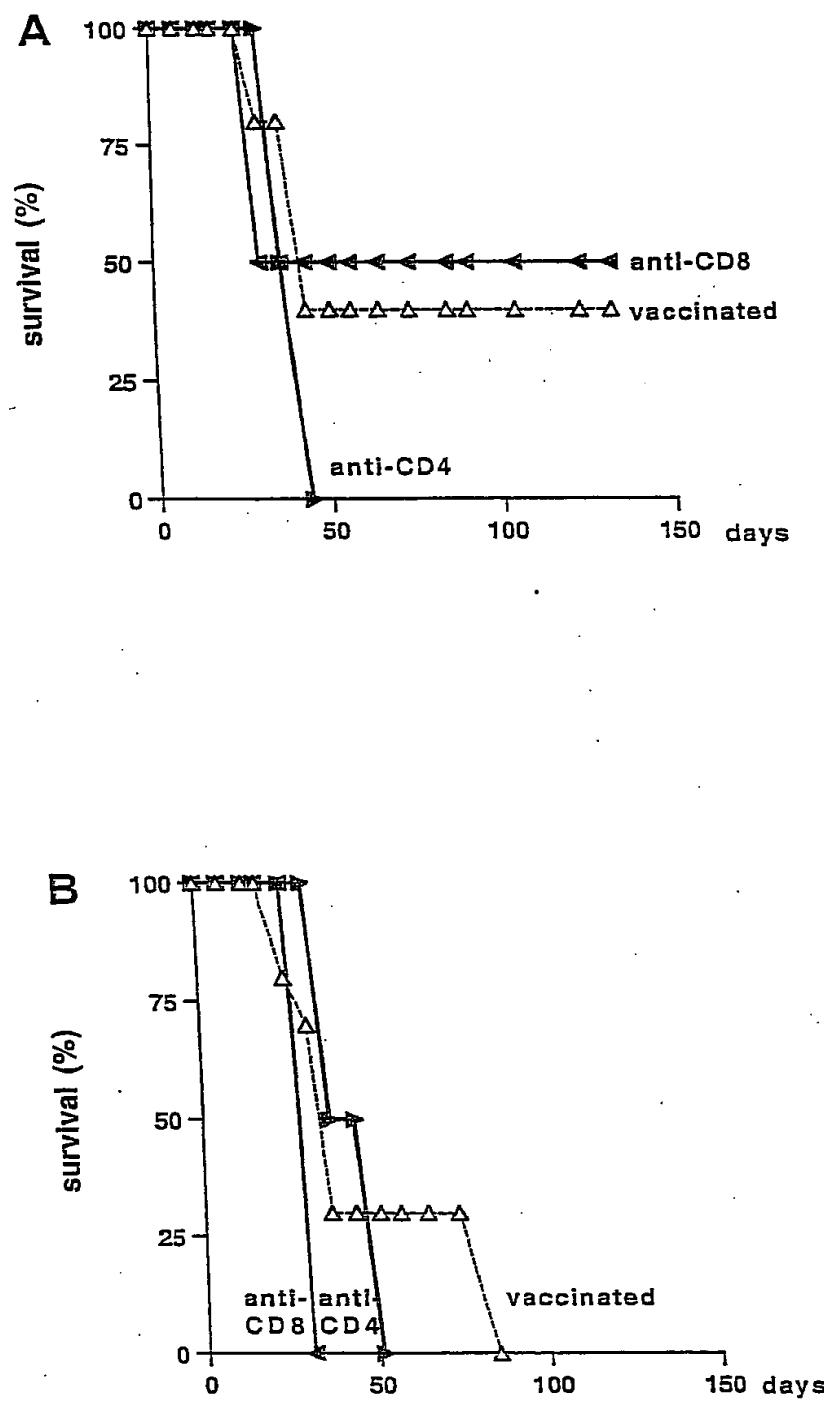


Fig. 3

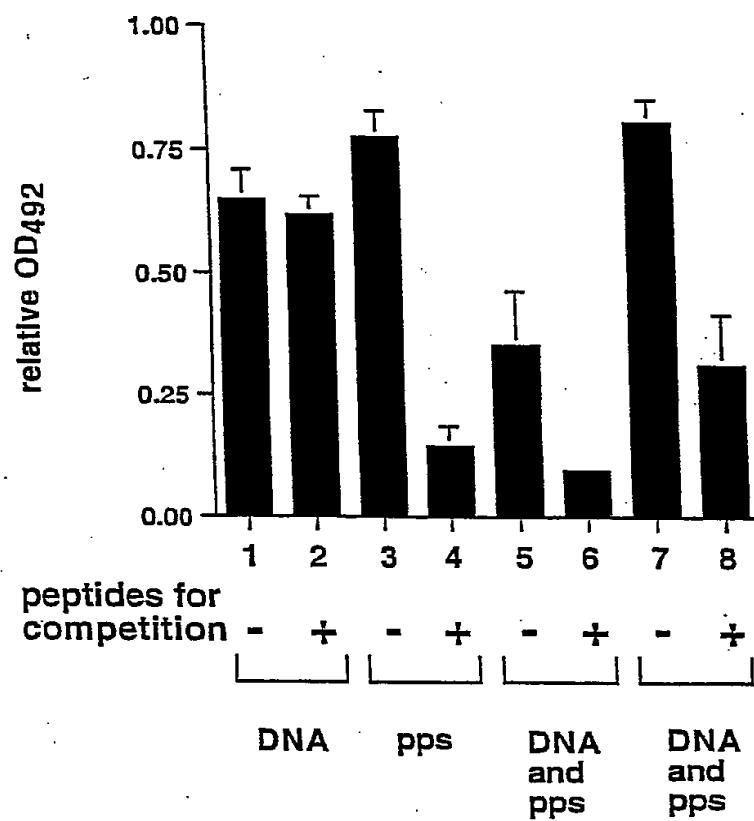


Fig. 4

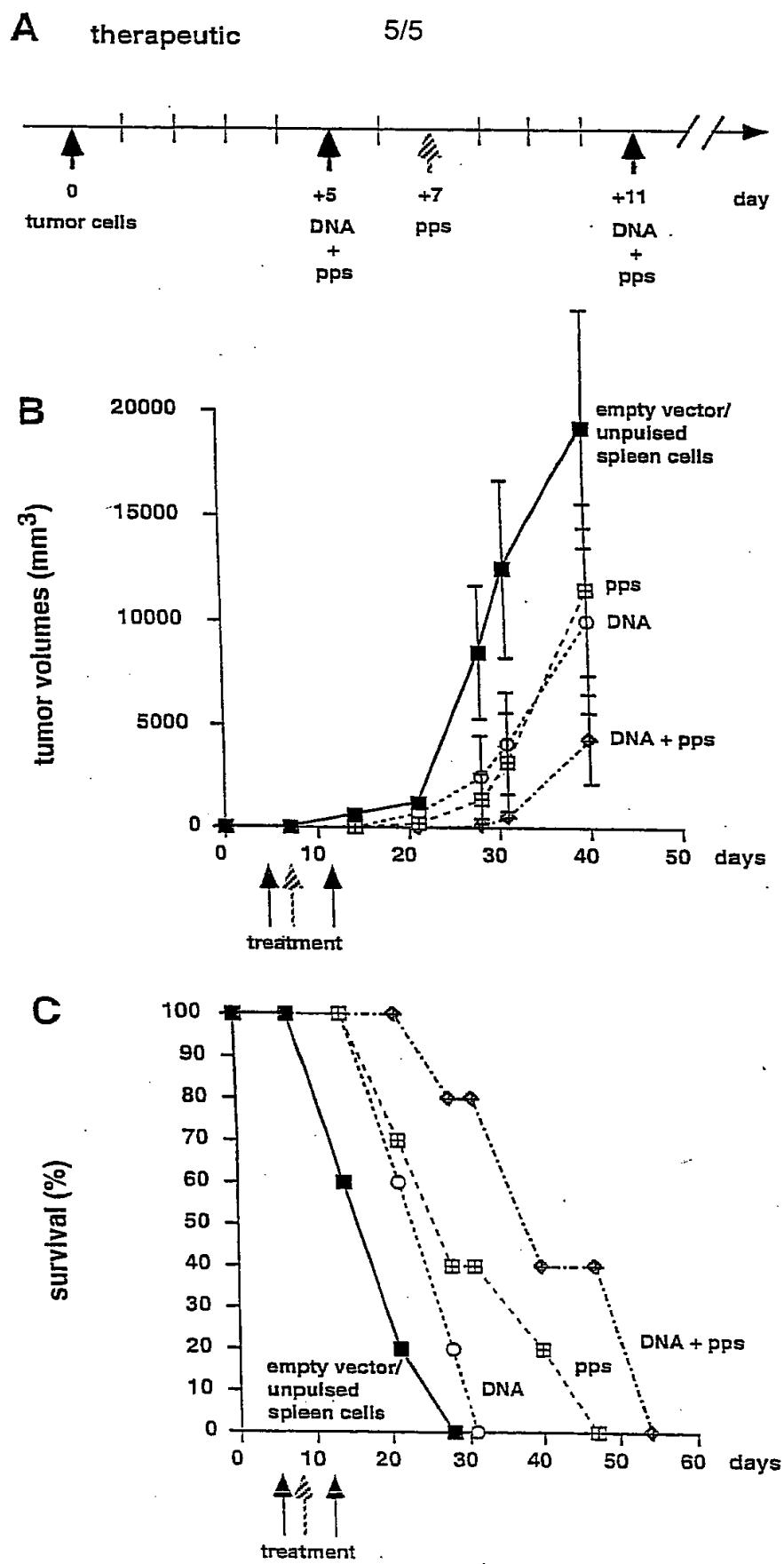


Fig. 5